

Dual Trade of Bcl-2 and Bcl-x_L in Islet Physiology

Balancing Life and Death With Metabolism Secretion Coupling

Bernat Soria^{1,2} and Benoit R. Gauthier¹

Apoptosis is now recognized as a predominant mechanism by which β cells are destroyed in both type 1 and type 2 diabetes (1). Pro- and antiapoptotic members of the Bcl-2 family are central regulators involved in β -cell fate decision between life and death in response to physiological insults. The intricate interplay and balance between members of this family regulate apoptosis by controlling mitochondrial cell death signaling, the so-called intrinsic pathway (2). The Bcl-2 family is divided into three subgroups as follows: 1) the antiapoptotic Bcl-2-like proteins (Bcl-2, Bcl-x_L, Bcl-W, Mcl-1, and A1/Bfl1) that possess four independent and highly conserved Bcl-2 homology (BH) domains (BH-1 to -4); 2) the proapoptotic Bax-like proteins (Bax, Bak, and Bok/Mtd), which also harbor the four BH domains; and 3) proapoptotic members typified by a single BH-3 domain (Bid, Bim/Bod, Bad, Bmf, Bik/Nbk, Blk, NoxA, Puma/Bbc3, and Hrk/DP5). Members of this last subgroup initiate the apoptotic-signaling cascade by interacting with antiapoptotic members (3). In addition, intracellular calcium activity ($[Ca^{2+}]_i$) is an important second messenger that initiates the apoptotic program (4) as well as being essential in metabolism secretion coupling that drives the Ca^{2+} sensor of the exocytotic machinery (5). Given the importance of Bcl-2 and Bcl-x_L as prosurvival sentinels of the mitochondria, combined with the indispensable function of this organelle in β -cell metabolism secretion coupling, these proteins have been the targets of numerous studies in which expression levels were manipulated with the goal of improving islet viability. The study by Zhou et al. (6) in 2000 was the first to provide proof of concept that overexpression of Bcl-x_L in β cells blunted stressed-induced apoptosis. Unexpectedly, one of the transgenic lines expressing high levels of Bcl-x_L also displayed severe hyperglycemia. Glucose intolerance in these animals was linked to impaired nutrient-induced insulin secretion caused by alterations in mitochondrial ATP generation and intracellular Ca^{2+} handling (6). Substantiating this study, Pax4-induced Bcl-x_L expression in rat islets also resulted in curtailed β -cell metabolism-secretion coupling through

blunted mitochondrial Ca^{2+} concentrations and decreased ATP production (7). These early studies cautioned that Bcl-2 protein family members might not only be involved in regulating apoptosis but also be implicated in a day-to-day control of cellular function. Consistent with this premise was the elegant finding by Danial et al. (8) that Bad forms a complex with glucokinase in order to regulate glucose-driven mitochondrial metabolism and insulin secretion in islets. Nonetheless, other studies have tended to argue against the role of these proteins in glucose metabolism. Indeed, transgenic mice either overexpressing Bcl-2 or bearing a deletion in the *BCLXL* gene specifically in β cells were not reported to display any aberrant alterations in glucose metabolism (9,10). Thus, the relative contributions of antiapoptotic versus alternative functions of Bcl-2 and Bcl-x_L for overall β -cell survival and performance remain controversial.

In this issue of *Diabetes*, Luciani et al. (11) address this controversy and convincingly demonstrate that Bcl-2 or Bcl-x_L dampens glucose-induced insulin secretion and highlight the role of these prosurvival proteins as critical physiological integrators balancing life and death with metabolism secretion coupling in the β cell. In a first approach to authenticate this dual functionality, the authors used the small-molecule antagonist compound 6 (C6) and YC137 to pharmacologically hinder Bcl-2 and Bcl-x_L. These antagonists bind to and displace proapoptotic members such as Bad from Bcl-2 and Bcl-x_L, ultimately inducing apoptosis. In these experiments, C6 caused a rapid disruption of the Bcl-x_L/Bad complex as well as a redistribution of Bax from the cytosol to mitochondria resulting in the release of cytochrome c, activation of caspase-3, and β -cell death. As antagonist-induced apoptosis was usually detected 2 h posttreatment, the authors argued that cellular events occurring within this time frame were likely independent of the central apoptotic events. In this context, the most impressive physiological event occurring subsequent to antagonistic treatment was the rapid triggering of $[Ca^{2+}]_i$ in cells that mimicked the effect of glucose signaling. Yet, cells were cultured in the presence of low glucose, suggesting increased performance of mitochondrial metabolism leading to Ca^{2+} influx and potentially insulin secretion. Luciani et al. (11) methodically dissect the pathway leading to glucose-induced insulin secretion using various inhibitors and demonstrate that antagonizing Bcl-2/Bcl-x_L in islets recapitulates cellular events associated with metabolism secretion coupling in β -cells: increased ATP production causing closure of the ATP-sensitive K^+ channel with the subsequent depolarization of the plasma membrane and opening of the L-type Ca^{2+} channel resulting in submembranous increase in $[Ca^{2+}]_i$ and ultimately insulin exocytosis. Low glucose levels as well as a sustained

From the ¹Department of Stem Cells, Andalusian Center for Molecular Biology and Regenerative Medicine, Seville, Spain; and the ²Centro de Investigación Biomédica en Red de Diabetes y Enfermedades Metabólicas Asociadas, Barcelona, Spain.

Corresponding author: Benoit R. Gauthier, benoit.gauthier@cabimer.es.
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Taken together, these data are noteworthy, as they provide the first convincing evidence that Bcl-2 and Bcl-x_L take on dual functions in β cells: on the one hand, they are the gatekeepers of life and death, and on the other they are the “thermostat” of energy production in mitochondria. In fact, we would like to propose the term “energystat” to describe this new regulatory function of Bcl-2 and Bcl-x_L. This is particularly relevant in a cell that lacks the Pasteur effect, a condition for being a nutrient sensor (12). In fact, these two roles are likely not mutually exclusive, as they converge on mitochondrial processes that will ultimately preserve β cells from deleterious stress. Indeed, as proposed by Luciani et al., restricting glucose metabolism may be a means by which the nonapoptotic function of Bcl-2 and Bcl-x_L protects β cells against reactive oxygen species generated through oxidative phosphorylation while the antiapoptotic function preserves mitochondrial integrity under metabolic stress conditions such as hyperglycemia. Interestingly, Bax, another



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member of the Bcl-2 family, was recently linked to mitochondrial energy production. Indeed, BAX-deficient HCT-116 colorectal cancer cells were shown to have blunted ATP biosynthesis, a metabolic alteration associated with reduced citrate synthase activity. In contrast, overexpression of Bcl-2 in wild-type HCT-116 cells caused a drastic decrease in ATP production. The authors of this study concluded that Bcl-2 impedes Bax action on mitochondrial bioenergetics by potentially blunting its interaction with other mitochondrial proteins (13). However, it is now clear from the study of Luciani et al. (11) that Bcl-2 suppresses glucose-driven mitochondrial ATP biosynthesis independently of Bax or Bak. Thus, Bcl-2 and Bcl-x_L join the armada of apoptotic factors along with Bad and Bax that possesses dual functionality. Despite these findings, one outstanding question remains to be clearly resolved from the study of Luciani et al.: what are the downstream mitochondrial targets mediating the nonapoptotic effects of Bcl-2 and Bcl-x_L on β -cell bioenergetics? Experimental data would tend to suggest that the answer to this question lies within the tricarboxylic acid (TCA) cycle and/or the oxidative phosphorylation metabolic pathways. Indeed, 1) levels of acetyl-CoA as well as those of citrate, α -ketoglutarate, and succinate, three key metabolites of the TCA cycle, were reduced in Bcl-x_L-overexpressing Jurkat cells (14); 2) pyruvate, the main substrate for the generation of mitochondrial acetyl-CoA that fuels the TCA cycle, was unable to rescue impaired insulin secretion in Bcl-x_L-overexpressing islets (6); and 3) Bcl-x_L was shown to interact with the ATP synthase β -subunit and to regulate mitochondrial energetics by stabilizing the inner membrane potential in neuronal cells (15). In addition, Bax was found to interact with citrate synthase, which is involved in the production of citrate from acetyl-CoA and oxaloacetate (16). Thus, it is tempting to speculate that Bcl-2 family members interact with and regulate the activity of key TCA cycle enzymes and proteins of the respiratory chain to energize levels of ATP biosynthesis (Fig. 1). Two potential candidate targets could be the pyruvate dehydrogenase complex or the pyruvate carboxylase. These control the entry of pyruvate in the form of acetyl-CoA and oxaloacetate, respectively, into the TCA cycle. Alternatively, Bcl-2 or Bcl-x_L could alter levels of acetyl-CoA by interacting with the citrate carrier, which is responsible for the efflux of this metabolite from the mitochondria to the cytosol in the form of citrate. Luciani et al. (11) also propose that the voltage-dependent anion channel and the mitochondrial adenine nucleotide translocator, which are the main venues by which ATP and ADP are exchanged between the mitochondrial matrix and the cytosol, could also be targets of Bcl-2 and Bcl-x_L.

While these various targets are plausible Bcl-2 and Bcl-x_L regulatory checkpoints, several issues remain to be addressed from the study of Luciani et al. in order to firmly validate the bona fide implication of these survival factors in the management of mitochondrial metabolism. Indeed, the antagonists C6 and YC137 were shown to preferentially bind to and inhibit Bcl-2 while displaying low affinity for Bcl-x_L. Consistent with the latter, islets derived from Bcl-2-ablated transgenic mice displayed a stronger precocious insulin secretion response to lower glucose concentrations compared with Bclx β KO islets (11). Thus, although rapid triggering of $[Ca^{2+}]_i$ was also observed in Bclx β KO islets, the direct impact of Bcl-x_L on insulin secretion needs to be

further investigated. Furthermore, as YC137 also inhibits Bcl-W, which is strongly expressed in islets, the role of this prosurvival factor in regulating mitochondrial bioenergetics needs to be addressed (17,18). This is of particular interest, as human islets express low levels of Bcl-2 relative to other family members such as Bcl-W (19).

In summary, the study by Luciani et al. raises serious concerns about the feasibility of using prosurvival members of the Bcl-2 family as therapeutic targets for the treatment of diabetes. Indeed, human islets cultured in high glucose were found to have reduced levels of Bcl-x_L (20). This most likely alleviates constraints on glucose-driven mitochondrial ATP production, thereby allowing increased insulin secretion to restore normal circulating glucose levels. Nonetheless, decreased Bcl-x_L levels also tilt the balance between pro- and antiapoptotic factors favoring apoptosis. Paradoxically, cell death may be instrumental in this context to induce β -cell renewal in order to maintain a healthy and functional β -cell mass. Indeed, a recent study demonstrated that insulinoma INS-1 cells undergoing caspase-dependent apoptosis promoted the regenerative capacity of neighboring cells by shedding microparticles harboring the pancreatic stone protein/regenerating protein (21). Thus, Bcl-2 protein family members clearly act as double-edged swords in the context of diabetes. Extreme care should be taken to establish whether increased or decreased expression of these proteins would provide the most favorable outcome for the treatment of diabetes.

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